



39780-1216R1D4 SAVED AUGUST 17 2005.TXT

SEQUENCE LISTING

<110> Ashkenazi, Avi J.  
Fong, Sherman  
Goddard, Audrey  
Gurney, Austin L.  
Napier, Mary A.  
Tumas, Daniel  
Wood, William I.

<120> COMPOUNDS, COMPOSITIONS AND METHODS FOR  
THE TREATMENT OF DISEASES CHARACTERIZED BY A33- RELATED  
ANTIGENS

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<151> 1999-03-05

<150> PCT/US98/24855

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 65 70 75 80  
 Pro Tyr Ala Asp Arg Val Thr Phe Ser Ser Gly Ile Thr Phe Ser  
 85 90 95  
 Ser Val Thr Arg Lys Asp Asn Gly Glu Tyr Thr Cys Met Val Ser Glu  
 100 105 110  
 Glu Gly Gly Gln Asn Tyr Gly Glu Val Ser Ile His Leu Thr Val Leu  
 115 120 125  
 Val Pro Pro Ser Lys Pro Thr Ile Ser Val Pro Ser Val Thr Ile  
 130 135 140  
 Gly Asn Arg Ala Val Leu Thr Cys Ser Glu His Asp Gly Ser Pro Pro  
 145 150 155 160  
 Ser Glu Tyr Ser Trp Phe Lys Asp Gly Ile Ser Met Leu Thr Ala Asp  
 165 170 175  
 Ala Lys Lys Thr Arg Ala Phe Met Asn Ser Ser Phe Thr Ile Asp Pro  
 180 185 190  
 Lys Ser Gly Asp Leu Ile Phe Asp Pro Val Thr Ala Phe Asp Ser Gly  
 195 200 205  
 Glu Tyr Tyr Cys Gln Ala Gln Asn Gly Tyr Gly Thr Ala Met Arg Ser  
 210 215 220  
 Glu Ala Ala His Met Asp Ala Val Glu Leu Asn Val Gly Gly Ile Val  
 225 230 235 240  
 Ala Ala Val Leu Val Thr Leu Ile Leu Leu Gly Leu Leu Ile Phe Gly  
 245 250 255  
 Val Trp Phe Ala Tyr Ser Arg Gly Tyr Phe Glu Thr Thr Lys Lys Gly  
 260 265 270  
 Thr Ala Pro Gly Lys Lys Val Ile Tyr Ser Gln Pro Ser Thr Arg Ser  
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 Glu Gly Glu Phe Lys Gln Thr Ser Ser Phe Leu Val  
 290 295 300

<210> 11  
 <211> 1842  
 <212> DNA  
 <213> Homo sapiens

<400> 11  
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 aaggcgcaag tcgagaggaa actgttgtgc ctcttcatat tggcgatcct gttgtgctcc 120  
 ctggcattgg gcagtgttac agtgcactct tctgaacctg aagtcagaat tcctgagaat 180  
 aatcctgtga agttgtcctg tgcctactcg ggcttttctt ctccccgtgt ggagtgggaag 240  
 tttgaccaag gagacaccac cagactcgtt tgctataata acaagatcac agcttcctat 300  
 gaggaccggg tgaccttctt gccaaactggt atcaccttca agtccgtgac acgggaagac 360  
 actgggacat acacttgtat ggtctctgag gaaggcggca acagctatgg ggaggtcaag 420  
 gtcaagctca tcgtgcttgt gcctccatcc aagcctacag ttaacatccc ctctctgcc 480  
 accattggga accgggcagt gctgacatgc tcagaacaag atggttcccc accttctgaa 540

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tacacctggt	tcaaagatgg	gatagtgatg	cctacgaatc	ccaaaagcac	ccgtgccttc	600
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gcctctgata	ctggagaata	cagctgtgag	gcacggaatg	ggatgaggac	acccatgact	720
tcaaattgctg	tgcgcatgga	agctgtggag	cggaatgtgg	gggtcatcgt	ggcagccgtc	780
cttgtaaccc	tgattctcct	gggaatcttg	gtttttggca	tctggtttgc	ctatagccga	840
ggccactttg	acagaacaaa	gaaagggact	tcgagtaaga	aggtgattta	cagccagcct	900
agtgtcccga	gtgaaggaga	attcaaacag	acctcgtcac	tcctgggtgtg	agcctggtcg	960
gctcaccgcc	tatcatctgc	atttgcctta	ctcaggtgct	accggactct	ggcccctgat	1020
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tcggatgtgt	ttttaataat	gtcagctatg	tgccccatcc	tccttcacgc	cctccctccc	1140
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agggatcagg	aaggaatcct	gggtatgcc	ttgacttccc	ttctaagtag	acagcaaaaa	1260
tggtcggggtg	cgcaggaatc	tgacttcaac	tgcccacctg	gctggaaggg	atctttgaat	1320
aggtatcttg	agcttggttc	tgggctcttt	ctttgtgtac	tgacgaccag	ggccagctgt	1380
tctagagcgg	gaattagagg	ctagagcggc	tgaatgggtt	gtttgggtgat	gacactgggg	1440
tccttccatc	tctggggccc	actctcttct	gtcttcccat	gggaagtgcc	actgggatcc	1500
ctctgccctg	tcctcctgaa	tacaagctga	ctgacattga	ctgtgtctgt	ggaaaatggg	1560
agctcttgtt	gtggagagca	tagtaaattt	tcagagaact	tgaagccaaa	aggattttaa	1620
accgctgtct	taaagaaaag	aaaactggag	gctgggcgca	gtggctcacg	cctgtaatcc	1680
cagaggctga	ggcaggcgga	tcacctgagg	tcgggagttc	gggatcagcc	tgaccaacat	1740
ggagaaaccc	tactggaaat	acaaagttag	ccaggcatgg	tggtgcatgc	ctgtagtccc	1800
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<210> 12  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Synthetic Oligonucleotide Primer

<400> 12  
 tcgctggagct gtgttctgtt tccc 24

<210> 13  
 <211> 50  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Synthetic Oligonucleotide Hybridization Probe

<400> 13  
 tgatcgcgat ggggacaaag gcgcaagctc gagaggaaac tggtgtgcct 50

<210> 14  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Synthetic Oligonucleotide Primer

<400> 14  
 acacctgggtt caaagatggg 20

<210> 15  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Synthetic Oligonucleotide Primer



<400> 15  
taggaagagt tgctgaaggc acgg 24

<210> 16  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic Oligonucleotide Primer

<400> 16  
ttgccttact caggtgctac 20

<210> 17  
<211> 20  
<212> DNA  
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<220>  
<223> Synthetic Oligonucleotide Primer

<400> 17  
actcagcagt ggtaggaaag 20

<210> 18  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic Oligonucleotide Primer

<400> 18  
tatccctcca attgagcacc ctgg 24

<210> 19  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic Oligonucleotide Primer

<400> 19  
gtcggaagac atcccaacaa g 21

<210> 20  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic Oligonucleotide Primer

<400> 20  
cttcacaatg tcgctgtgct gctc 24

<210> 21  
<211> 24  
<212> DNA  
<213> Artificial Sequence

&lt;220&gt;

&lt;223&gt; Synthetic Oligonucleotide Primer

&lt;400&gt; 21

agccaaatcc agcagctggc ttac

24

&lt;210&gt; 22

&lt;211&gt; 50

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Synthetic Oligonucleotide Hybridization Probe

&lt;400&gt; 22

tggatgaccg gagccactac acgtgtgaag tcacctggca gactcctgat

50

&lt;210&gt; 23

&lt;211&gt; 260

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 23

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Leu Ala Leu Gly Ser Val Thr Val His Ser Ser Glu Pro Glu Val Arg
 1      5      10      15
Ile Pro Glu Asn Asn Pro Val Lys Leu Ser Cys Ala Tyr Ser Gly Phe
      20      25      30
Ser Ser Pro Arg Val Glu Trp Lys Phe Asp Gln Gly Asp Thr Thr Arg
      35      40      45
Leu Val Cys Tyr Asn Asn Lys Ile Thr Ala Ser Tyr Glu Asp Arg Val
      50      55      60
Thr Phe Leu Pro Thr Gly Ile Thr Phe Lys Ser Val Thr Arg Glu Asp
      65      70      75      80
Thr Gly Thr Tyr Thr Cys Met Val Ser Glu Glu Gly Gly Asn Ser Tyr
      85      90      95
Gly Glu Val Lys Val Lys Leu Ile Val Leu Val Pro Pro Ser Lys Pro
      100      105      110
Thr Val Asn Ile Pro Ser Ser Ala Thr Ile Gly Asn Arg Ala Val Leu
      115      120      125
Thr Cys Ser Glu Gln Asp Gly Ser Pro Pro Ser Glu Tyr Thr Trp Phe
      130      135      140
Lys Asp Gly Ile Val Met Pro Thr Asn Pro Lys Ser Thr Arg Ala Phe
      145      150      155      160
Ser Asn Ser Ser Tyr Val Leu Asn Pro Thr Thr Gly Glu Leu Val Phe
      165      170      175
Asp Pro Leu Ser Ala Ser Asp Thr Gly Glu Tyr Ser Cys Glu Ala Arg
      180      185      190
Asn Gly Tyr Gly Thr Pro Met Thr Ser Asn Ala Val Arg Met Glu Ala
      195      200      205
Val Glu Arg Asn Val Gly Val Ile Val Ala Ala Val Leu Val Thr Leu
      210      215      220
Ile Leu Leu Gly Ile Leu Val Phe Gly Ile Trp Phe Ala Tyr Ser Arg
      225      230      235      240
Gly His Phe Asp Arg Thr Lys Lys Gly Thr Ser Ser Lys Lys Val Ile
      245      250      255
Tyr Ser Gln Pro
      260

```

&lt;210&gt; 24

&lt;211&gt; 270

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 24

```

Val Arg Val Thr Val Asp Ala Ile Ser Val Glu Thr Pro Gln Asp Val
1      5      10      15
Leu Arg Ala Ser Gln Gly Lys Ser Val Thr Leu Pro Cys Thr Tyr His
20      25      30
Thr Ser Thr Ser Ser Arg Glu Gly Leu Ile Gln Trp Asp Lys Leu Leu
35      40      45
Leu Thr His Thr Glu Arg Val Val Ile Trp Pro Phe Ser Asn Lys Asn
50      55      60
Tyr Ile His Gly Glu Leu Tyr Lys Asn Arg Val Ser Ile Ser Asn Asn
65      70      75      80
Ala Glu Gln Ser Asp Ala Ser Ile Thr Ile Asp Gln Leu Thr Met Ala
85      90      95
Asp Asn Gly Thr Tyr Glu Cys Ser Val Ser Leu Met Ser Asp Leu Glu
100     105     110
Gly Asn Thr Lys Ser Arg Val Arg Leu Leu Val Leu Val Pro Pro Ser
115     120     125
Lys Pro Glu Cys Gly Ile Glu Gly Glu Thr Ile Ile Gly Asn Asn Ile
130     135     140
Gln Leu Thr Cys Gln Ser Lys Glu Gly Ser Pro Thr Pro Gln Tyr Ser
145     150     155     160
Trp Lys Arg Tyr Asn Ile Leu Asn Gln Glu Gln Pro Leu Ala Gln Pro
165     170     175
Ala Ser Gly Gln Pro Val Ser Leu Lys Asn Ile Ser Thr Asp Thr Ser
180     185     190
Gly Tyr Tyr Ile Cys Thr Ser Ser Asn Glu Glu Gly Thr Gln Phe Cys
195     200     205
Asn Ile Thr Val Ala Val Arg Ser Pro Ser Met Asn Val Ala Leu Tyr
210     215     220
Val Gly Ile Ala Val Gly Val Val Ala Ala Leu Ile Ile Ile Gly Ile
225     230     235     240
Ile Ile Tyr Cys Cys Cys Cys Arg Gly Lys Asp Asp Asn Thr Glu Asp
245     250     255
Lys Glu Asp Ala Arg Pro Asn Arg Glu Ala Tyr Glu Glu Pro
260     265     270

```

&lt;210&gt; 25

&lt;211&gt; 263

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 25

```

Leu Cys Ser Leu Ala Leu Gly Ser Val Thr Val His Ser Ser Glu Pro
1      5      10      15
Glu Val Arg Ile Pro Glu Asn Asn Pro Val Lys Leu Ser Cys Ala Tyr
20      25      30
Ser Gly Phe Ser Ser Pro Arg Val Glu Trp Lys Phe Asp Gln Gly Asp
35      40      45
Thr Thr Arg Leu Val Cys Tyr Asn Asn Lys Ile Thr Ala Ser Tyr Glu
50      55      60
Asp Arg Val Thr Phe Leu Pro Thr Gly Ile Thr Phe Lys Ser Val Thr
65      70      75      80
Arg Glu Asp Thr Gly Thr Tyr Thr Cys Met Val Ser Glu Glu Gly Gly
85      90      95
Asn Ser Tyr Gly Glu Val Lys Val Lys Leu Ile Val Leu Val Pro Pro
100     105     110
Ser Lys Pro Thr Val Asn Ile Pro Ser Ser Ala Thr Ile Gly Asn Arg
115     120     125
Ala Val Leu Thr Cys Ser Glu Gln Asp Gly Ser Pro Pro Ser Glu Tyr
130     135     140

```

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```

Thr Trp Phe Lys Asp Gly Ile Val Met Pro Thr Asn Pro Lys Ser Thr
145      150      155      160
Arg Ala Phe Ser Asn Ser Ser Tyr Val Leu Asn Pro Thr Thr Gly Glu
      165      170      175
Leu Val Phe Asp Pro Leu Ser Ala Ser Asp Thr Gly Glu Tyr Ser Cys
      180      185      190
Glu Ala Arg Asn Gly Tyr Gly Thr Pro Met Thr Ser Asn Ala Val Arg
      195      200      205
Met Glu Ala Val Glu Arg Asn Val Gly Val Ile Val Ala Ala Val Leu
      210      215      220
Val Thr Leu Ile Leu Leu Gly Ile Leu Val Phe Gly Ile Trp Phe Ala
225      230      235      240
Tyr Ser Arg Gly His Phe Asp Arg Thr Lys Lys Gly Thr Ser Ser Lys
      245      250      255
Lys Val Ile Tyr Ser Gln Pro
      260

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<210> 26  
 <211> 273  
 <212> PRT  
 <213> Homo sapiens

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<400> 26
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Gln Asp Val Leu Arg Ala Ser Gln Gly Lys Ser Val Thr Leu Pro Cys
      20      25      30
Thr Tyr His Thr Ser Thr Ser Arg Glu Gly Leu Ile Gln Trp Asp
      35      40      45
Lys Leu Leu Thr His Thr Glu Arg Val Val Ile Trp Pro Phe Ser
      50      55      60
Asn Lys Asn Tyr Ile His Gly Glu Leu Tyr Lys Asn Arg Val Ser Ile
65      70      75      80
Ser Asn Asn Ala Glu Gln Ser Asp Ala Ser Ile Thr Ile Asp Gln Leu
      85      90      95
Thr Met Ala Asp Asn Gly Thr Tyr Glu Cys Ser Val Ser Leu Met Ser
      100      105      110
Asp Leu Glu Gly Asn Thr Lys Ser Arg Val Arg Leu Leu Val Leu Val
      115      120      125
Pro Pro Ser Lys Pro Glu Cys Gly Ile Glu Gly Glu Thr Ile Ile Gly
      130      135      140
Asn Asn Ile Gln Leu Thr Cys Gln Ser Lys Glu Gly Ser Pro Thr Pro
145      150      155      160
Gln Tyr Ser Trp Lys Arg Tyr Asn Ile Leu Asn Gln Glu Gln Pro Leu
      165      170      175
Ala Gln Pro Ala Ser Gly Gln Pro Val Ser Leu Lys Asn Ile Ser Thr
      180      185      190
Asp Thr Ser Gly Tyr Tyr Ile Cys Thr Ser Ser Asn Glu Glu Gly Thr
      195      200      205
Gln Phe Cys Asn Ile Thr Val Ala Val Arg Ser Pro Ser Met Asn Val
      210      215      220
Ala Leu Tyr Val Gly Ile Ala Val Gly Val Val Ala Ala Leu Ile Ile
225      230      235      240
Ile Gly Ile Ile Ile Tyr Cys Cys Cys Cys Arg Gly Lys Asp Asp Asn
      245      250      255
Thr Glu Asp Lys Glu Asp Ala Arg Pro Asn Arg Glu Ala Tyr Glu Glu
      260      265      270
Pro

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<210> 27

<211> 413

<212> DNA

<213> Artificial Sequence

<220>

<223> Consensus DNA Sequence

<400> 27

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aggccaaaac ctggaagagg atacagtcac tctggaagta ttagtggtc cagcagttcc 120
atcatgtgaa gtaccctctt ctgctctgag tggaaactgtg gtagagctac gatgtcaaga 180
caaagaaggg aatccagctc ctgaatacac atggtttaag gatggcatcc gtttgctaga 240
aaatcccaga cttggctccc aaagcaccaa cagctcatal acaatgaata caaaaactgg 300
aactctgcaa ttttaatactg tttccaaact ggacactgga gaatattcct gtgaagcccg 360
caattctgtt ggatatcgca ggtgtcctgg ggaaacgaat gcaagtagat gat 413
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<210> 28

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Primer

<400> 28

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atcgttgtga agttagtgcc cc 22
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<210> 29

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Primer

<400> 29

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acctgcgata tccaacagaa ttg 23
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<210> 30

<211> 48

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Hybridization Probe

<400> 30

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ggaagaggat acagtcactc tggaagtatt agtggctcca gcagttcc 48
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